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Sequence Listing was accepted.

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Reviewer: markspencer

Timestamp: [year=2008; month=6; day=12; hr=15; min=57; sec=14; ms=205;]

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Application No: 10506925 Version No: 3.0

Input Set:

Output Set:

Started: 2008-05-19 14:11:08.608
Finished: 2008-05-19 14:11:13.865
Elapsed: 0 hr(s) 0 min(s) 5 sec(s) 257 ms
Total Warnings: 17
Total Errors: 0
No. of SeqIDs Defined: 17
Actual SeqID Count: 17

Error code	Error Description
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<110> Neurogenex Co., Ltd.

<120> ENHANCED INSERTED YELLOW FLUORESCENCE PROTEIN AND ITS

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<141> 2004-09-07

<150> KR10-2002-0012409

<151> 2002-03-08

<150> KR10-2002-0015217

<151> 2002-03-21

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<151> 2002-03-21

<160> 17

<170> PatentIn version 3.4

<210> 1

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<212> PRT

<213> Artificial Sequence

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<223> y-citrine of fluorescence protein

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Val Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly
20 25 30

Glu Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile
35 40 45

Cys Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr
50 55 60

Phe Gly Tyr Gly Leu Met Cys Phe Ala Arg Tyr Pro Asp His Met Lys
65 70 75 80

Gln His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu
85 90 95

Arg Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu
100 105 110

Val Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly
115 120 125

Ile Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr
130 135 140

Asn Tyr Gly Gly Ser Gly Ala Ser Asn Ser His Asn Val Tyr Ile Met
145 150 155 160

Ala Asp Lys Gln Lys Asn Gly Ile Lys Val Asn Phe Lys Ile Arg His
165 170 175

Asn Ile Glu Asp Gly Ser Val Gln Leu Ala Asp His Tyr Gln Gln Asn
180 185 190

Thr Pro Ile Gly Asp Gly Pro Val Leu Leu Pro Asp Asn His Tyr Leu
195 200 205

Ser Tyr Gln Ser Ala Leu Ser Lys Asp Pro Asn Glu Lys Arg Asp His
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Met Val Leu Leu Glu Phe Val Thr Ala Ala Gly Ile Thr Leu Gly Met
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Asp Glu Leu Tyr Lys
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<211> 245

<212> PRT

<213> Artificial Sequence

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<223> Peridot of fluorescence protein

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Val Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly

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Glu Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile
 35 40 45

Cys Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr
 50 55 60

Phe Gly Tyr Gly Leu Met Cys Phe Ala Arg Tyr Pro Asp His Met Lys
 65 70 75 80

Gln His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu
 85 90 95

Arg Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu
 100 105 110

Val Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly
 115 120 125

Ile Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr
 130 135 140

Asn Tyr Gly Gly Ser Gly Ala Ser Asn Ser His Asn Val Tyr Ile Met
 145 150 155 160

Ala Asp Lys Gln Lys Asn Gly Ile Lys Val Asn Phe Lys Ile Arg His
 165 170 175

Asn Ile Glu Asp Gly Ser Val Gln Leu Ala Asp His Tyr Gln Gln Asn
 180 185 190

Thr Pro Ile Gly Asp Gly Leu Val Leu Leu Pro Asp Asn His Tyr Leu
 195 200 205

Ser Tyr Gln Ser Ala Leu Ser Lys Asp Pro Asn Glu Lys Arg Asp His
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Met Val Leu Leu Glu Phe Val Thr Ala Ala Gly Ile Thr Ile Gly Met
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Asp Glu Leu Tyr Lys
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33

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<212> DNA

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ggcaagctga ccttgaagtt catctgcacc accggcaagc tgcccgtgcc ctggcccacc 180

ctcgtgacta ctttcggcta cggcctgatg tgcttcgccc gctaccccgga ccacatgaag 240

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gtgaaccgca tcgagctgaa gggcatcgac ttcaaggagg acggcaacat cctggggcac 420

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gaatttaaag aggctttctc cctatttgac aaggacgggg atgggacaat aacaaccaag 540

gagctgggga cggatgatgc gtctctgggg cagaaccca cagaagcaga gctgcaggac 600

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atgatggcaa gaaaaatgaa agacacagac agtgaagaag aaattagaga agcgttccgt 720

gtgtttgata aggatggcaa tggtacatc agtgcagcag agcttcgcca cgtgatgaca 780

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aagatccgcc acaacatcga ggacggcagc gtgcagctcg ccgaccacta ccagcagaac 1020

accccatcg gcgacggcct cgtgctgctg cccgacaacc actacctgag ctaccagtcc 1080

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 <223> NheI/DEVD R primer

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1